

## SPECIFICATION AMENDMENT

Please amend the title of the application to read:

~~Efficiency~~Efficient Agrobacterium-Mediated Wheat Transformation Method

Please amend paragraph [0052] to read:

The abbreviations in the table are described as follows: The CaMV.35S:en (e35S) promoter is a modification of the 35S promoter derived from the 35S RNA of cauliflower mosaic virus (CaMV) which contains a duplication of the -90 to -300 region; The Os.Act1 promoter is from the rice actin 1 gene, including the promoter, the 5' UTR, and the first intron. The SCBV promoter is from sugarcane badnavirus and also includes the first rice actin intron (I-Os.act1). The L-Ta.Lha1 is the leader sequence from the wheat CAB gene. The GUS gene is the  $\beta$ -glucuronidase coding sequence ~~form~~from **E. coli**; The **nptII** gene codes for neomycin phosphotransferase; The CP4 gene codes for glyphosate resistance; the T.Agrtu.nos (nos3' region contains downstream untranslated sequence and the poly A signal for the NOS gene of **Agrobacterium tumefaciens** pTiT37; and the T-Ta.Hsp17 is the 3' region of the wheat heat shock protein 17. All elements are named using current conventions in the art as found at <http://genome-www.stanford.edu/mendel>.